



09/856723

SEQUENCE LISTING

<110> Kramer, Michael

<120> Regulatory Protein pKe#83 from Human
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<140> PCT/DE 99/03732

<141> 1999-11-19

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<151> 1998-11-26

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<170> PatentIn Ver. 2.1

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Arg Ser Leu Glu Cys Arg Ser Asp Pro Glu Ser Pro Ile Lys Lys Thr
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Ser Leu Ser Pro Thr Ser Lys Leu Gly Tyr Ser Tyr Ser Arg Asp Leu
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Asp Leu Ala Lys Lys His Ala Ser Leu Arg Gln Thr Glu Ser Asp
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Pro Asp Ala Asp Arg Thr Thr Leu Asn His Ala Asp His Ser Ser Lys
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Ile Val Gln His Arg Leu Leu Ser Arg Gln Glu Glu Leu Lys Glu Arg
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Ala Arg Val Leu Leu Glu Gln Ala Arg Arg Asp Ala Ala Leu Lys Ala
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Leu Pro Ser Tyr Gly Glu Met Ala Ala Glu Lys Leu Lys Glu Arg Ser
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Lys Ala Ser Gly Asp Glu Asn Asp Asn Ile Glu Ile Asp Thr Asn Glu
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Ala Leu Val Glu Lys Arg Leu Arg Tyr Leu Met Asp Thr Gly Arg Asn
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Thr Glu Glu Glu Ala Met Met Gln Glu Trp Phe Met Leu Val Asn
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20 25 30

Arg Ser Leu Glu Cys Arg Ser Asp Pro Glu Ser Pro Ile Lys Lys Thr
35 40 45

Ser Leu Ser Pro Thr Ser Lys Leu Gly Tyr Ser Tyr Ser Arg Asp Leu
50 55 60

Asp Leu Ala Lys Lys His Ala Ser Leu Arg Gln Thr Glu Ser Asp
65 70 75 80

Pro Asp Ala Asp Arg Thr Thr Leu Asn His Ala Asp His Ser Ser Lys
85 90 95

Ile Val Gln His Arg Leu Leu Ser Arg Gln Glu Leu Lys Glu Arg
100 105 110

Ala Arg Val Leu Leu Glu Gln Ala Arg Arg Asp Ala Ala Lys Ala
115 120 125

Gly Asn Lys His Asn Thr Asn Thr Ala Thr Pro Phe Cys Asn Arg Gln
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Leu Ser Asp Gln Gln Asp Glu Glu Arg Arg Arg Gln Leu Arg Glu Arg
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Asn Ser Lys Leu Val Asp Leu Lys Leu Lys Lys Leu Leu Glu Val Gln
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Pro Gln Val Ala Asn Ser Pro Ser Ser Ala Ala Gln Lys Ala Val Thr
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Arg Ser Leu Glu Cys Arg Ser Asp Pro Glu Ser Pro Ile Lys Lys Thr
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Ser Leu Ser Pro Thr Ser Lys Leu Gly Tyr Ser Tyr Ser Arg Asp Leu
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Asp Leu Ala Lys Lys His Ala Ser Leu Arg Gln Thr Glu Ser Asp
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Pro Asp Ala Asp Arg Thr Thr Leu Asn His Ala Asp His Ser Ser Lys
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Ile Val Gln His Arg Leu Leu Ser Arg Gln Glu Glu Leu Lys Glu Arg
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Ala Arg Val Leu Leu Glu Gln Ala Arg Arg Asp Ala Ala Leu Lys Ala
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Casein kinase, 5x Tyrosine kinase

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35 40 45

Met Gln Ser Leu Ala Ser Leu Val Ser Met Lys Gln Ala Asp Ile Gly
50 55 60

Asn Leu Asp Asp Phe Glu Glu Asp Asn Glu Asp Asp Asp Glu Asn Arg
65 70 75 80

Val Asn Gln Glu Glu Lys Ala Ala Lys Ile Thr Glu Leu Ile Asn Lys
85 90 95

Leu Asn Phe Leu Asp Glu Ala Glu Lys Asp Leu Ala Thr Val Asn Ser
100 105 110

Asn Pro Phe Asp Asp Pro Asp Ala Ala Glu Leu Asn Pro Phe Gly Asp
115 120 125

Pro Asp Ser Glu Glu Pro Ile Thr Glu Thr Ala Ser Pro Arg Lys Thr
130 135 140

Glu Asp Ser Phe Tyr Asn Asn Ser Tyr Asn Pro Phe Lys Glu Val Gln
145 150 155 160

Thr Pro Gln Tyr Leu Asn Pro Phe Asp Glu Pro Glu Ala Phe Val Thr
165 170 175

Ile Lys Asp Ser Pro Pro Gln Ser Thr Lys Arg Lys Asn Ile Arg Pro
180 185 190

Val Asp Met Ser Lys Tyr Leu Tyr Ala Asp Ser Ser Lys Thr Glu Glu
195 200 205

Glu Glu Leu Asp Glu Ser Asn Pro Phe Tyr Glu Pro Lys Ser Thr Pro
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Pro Pro Asn Asn Leu Val Asn Pro Val Gln Glu Leu Glu Thr Glu Arg
225 230 235 240

Arg Val Lys Arg Lys Ala Pro Ala Pro Pro Val Leu Ser Pro Lys Thr
245 250 255

Gly Val Leu Asn Glu Asn Thr Val Ser Ala Gly Lys Asp Leu Ser Thr
260 265 270

Ser Pro Lys Pro Ser Pro Ile Pro Ser Pro Val Leu Gly Arg Lys Pro
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Tyr Arg Gly Val Lys Ile Thr Asn Phe Thr Thr Ser Trp Arg Asn Gly
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Leu Ser Phe Cys Ala Ile Leu His His Phe Arg Pro Asp Leu Ile Asp
325 330 335

Tyr Lys Ser Leu Asn Pro Gln Asp Ile Lys Glu Asn Asn Lys Lys Ala
340 345 350

Tyr Asp Gly Phe Ala Ser Ile Gly Ile Ser Arg Leu Leu Glu Pro Ser
355 360 365

Asp Met Val Leu Leu Ala Ile Pro Asp Lys Leu Thr Val Met Thr Tyr
370 375 380

Leu Tyr Gln Ile Arg Ala His Phe Ser Gly Gln Glu Leu Asn Val Val
385 390 395 400

Gln Ile Glu Glu Asn Ser Ser Lys Ser Thr Tyr Lys Val Gly Asn Tyr
405 410 415

Glu Thr Asp Thr Asn Ser Ser Val Asp Gln Glu Lys Phe Tyr Ala Glu
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Leu Ser Asp Leu Lys Arg Glu Pro Glu Leu Gln Gln Pro Ile Ser Gly
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Ala Val Asp Phe Leu Ser Gln Asp Asp Ser Val Phe Val Asn Asp Ser
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Gly Val Gly Glu Ser Glu Ser Glu His Gln Thr Pro Asp Asp His Leu
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Ser Pro Ser Thr Ala Ser Pro Tyr Cys Arg Arg Thr Lys Ser Asp Thr
485 490 495

Glu Pro Gln Lys Ser Gln Gln Ser Ser Gly Arg Thr Ser Gly Ser Asp
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Asp Pro Gly Ile Cys Ser Asn Thr Asp Ser Thr Gln Ala Gln Val Leu
515 520 525

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530 535 540

Leu Tyr Val Ser Asp Lys Lys Lys Asp Met Ser Pro Pro Phe Ile Cys
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565 570 575

Leu Glu Lys Glu Lys Leu Glu Asn Ser Arg Ser Leu Glu Cys Arg Ser
580 585 590

Asp Pro Glu Ser Pro Ile Lys Lys Thr Ser Leu Ser Pro Thr Ser Lys
595 600 605

Leu Gly Tyr Ser Tyr Ser Arg Asp Leu Asp Leu Ala Lys Lys Lys His
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Ala Ser Leu Arg Gln Thr Glu Ser Asp Pro Asp Ala Asp Arg Thr Thr
625 630 635 640

Leu Asn His Ala Asp His Ser Ser Lys Ile Val Gln His Arg Leu Leu
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Ser Arg Gln Glu Glu Leu Lys Glu Arg Ala Arg Val Leu Leu Glu Gln
660 665 670

Ala Arg Arg Asp Ala Ala Leu Lys Ala Gly Asn Lys His Asn Thr Asn
675 680 685

Thr Ala Thr Pro Phe Cys Asn Arg Gln Leu Ser Asp Gln Gln Asp Glu
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Glu Arg Arg Arg Gln Leu Arg Glu Arg Ala Arg Gln Leu Ile Ala Glu
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Ala Arg Ser Gly Val Lys Met Ser Glu Leu Pro Ser Tyr Gly Glu Met
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Ala Ala Glu Lys Leu Lys Glu Arg Ser Lys Ala Ser Gly Asp Glu Asn
740 745 750

Asp Asn Ile Glu Ile Asp Thr Asn Glu Glu Ile Pro Glu Gly Phe Val
755 760 765

Val Gly Gly Asp Glu Leu Thr Asn Leu Glu Asn Asp Leu Asp Thr
770 775 780

Pro Glu Gln Asn Ser Lys Leu Val Asp Leu Lys Leu Lys Lys Leu Leu
785 790 795 800

Glu Val Gln Pro Gln Val Ala Asn Ser Pro Ser Ser Ala Ala Gln Lys
805 810 815

Ala Val Thr Glu Ser Ser Glu Gln Asp Met Lys Ser Gly Thr Glu Asp
820 825 830

Leu Arg Thr Glu Arg Leu Gln Lys Thr Thr Glu Arg Phe Arg Asn Pro
835 840 845

Val Val Phe Ser Lys Asp Ser Thr Val Arg Lys Thr Gln Leu Gln Ser
850 855 860

Phe Ser Gln Tyr Ile Glu Asn Arg Pro Glu Met Lys Arg Gln Arg Ser
865 870 875 880

Ile Gln Glu Asp Thr Lys Lys Gly Asn Glu Glu Lys Ala Ala Ile Thr
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Glu Thr Gln Arg Lys Pro Ser Glu Asp Glu Val Leu Asn Lys Gly Phe
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Lys Asp Thr Ser Gln Tyr Val Val Gly Glu Leu Ala Ala Leu Glu Asn
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Arg Tyr Glu Leu Leu Asn Arg Glu Leu Arg Ala Met Leu Ala Ile Glu
995 1000 1005

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